

# REPLACEMENT SHEET

23132	CFR-1	MG160	1	MAACGRVRRM	FRLSAALHLL	LLFAAGGRNS	PARASHSOGQ	GPGANFVSFV	GQAGGGGPAG	QQLPQLPOSS	QLOQQQQQQQ	QOQQPQPPQ	PFPAAGPP-R
			1	MAACGRVRRM	FRLSAALHLL	LLFAAGAEKL	PGHGVHSOGQ	GPGANFVSFV	GQAGGGGPAG	QQLPQLPOSS	QLOQQQQQQQ	QOQQPQPPQ	PFPAAGPPAR
23132	CFR-1	MG160	101	RGAGAGGGW	KLAEEESCRE	DVTRVCPKHT	WSNNLAVIEC	QDVREPENE	ISSDCNHLLW	NYKLNLTDDP	KFESVAREVC	KSTITEIEEC	ADEPVGKGYM
			101	RGAGAGGGW	KLAEEESCRE	DVTRVCPKHT	WSNNLAVIEC	QDVREPENE	ISSDCNHLLW	NYKLNLTDDP	KFESVAREVC	KSTITEIEEC	ADEPVGKGYM
			101	RGAGAGGGW	KLAEEESCRE	DVTRVCPKHT	WSNNLAVIEC	QDVREPENE	ISSDCNHLLW	NYKLNLTDDP	KFESVAREVC	KSTITEIEEC	ADEPVGKGYM
23132	CFR-1	MG160	201	VSCLDVHRGN	ITEYQCHQYI	TKMTAIFSD	YRLICGFMDD	CKNDINILK	GSIRLGEKDA	HSQGEVVSCL	EKGLVKEAEE	REPKIQVSEL	CKKAILRVAE
			201	VSCLDVHRGN	ITEYQCHQYI	TKMTAIFSD	YRLICGFMDD	CKNDINILK	GSIRLGEKDA	HSQGEVVSCL	EKGLVKEAEE	REPKIQVSEL	CKKAILRVAE
			201	VSCLDVHRGN	ITEYQCHQYI	TKMTAIFSD	YRLICGFMDD	CKNDINILK	GSIRLGEKDA	HSQGEVVSCL	EKGLVKEAEE	REPKIQVSEL	CKKAILRVAE
23132	CFR-1	MG160	301	LSSDDPHLDR	HLFYACRDDR	ERFCENTQAG	EGRVYKCLFN	HKFEESMSEK	CREALTTROK	LIAQDYKVSY	SLAKSCKSDL	KKYRCNVENL	PRGREARLSY
			301	LSSDDPHLDR	HLFYACRDDR	ERFCENTQAG	EGRVYKCLFN	HKFEESMSEK	CREALTTROK	LIAQDYKVSY	SLAKSCKSDL	KKYRCNVENL	PRGREARLSY
			301	LSSDDPHLDR	HLFYACRDDR	ERFCENTQAG	EGRVYKCLFN	HKFEESMSEK	CREALTTROK	LIAQDYKVSY	SLAKSCKSDL	KKYRCNVENL	PRGREARLSY
23132	CFR-1	MG160	401	LLMCLES AVH	RGQVSSECO	GEMLDYRRML	MEDFSLSP EI	ILSCRGELIEH	HCSGLHRKGR	TLHCLMKVVR	GEKGNLGMNC	QOALQTLIQE	TDFGADYRID
			401	LLMCLES AVH	RGQVSSECO	GEMLDYRRML	MEDFSLSP EI	ILSCRGELIEH	HCSGLHRKGR	TLHCLMKVVR	GEKGNLGMNC	QOALQTLIQE	TDFGADYRID
			401	LLMCLES AVH	RGQVSSECO	GEMLDYRRML	MEDFSLSP EI	ILSCRGELIEH	HCSGLHRKGR	TLHCLMKVVR	GEKGNLGMNC	QOALQTLIQE	TDFGADYRID
23132	CFR-1	MG160	501	RALNEACESV	IQTACKHIRS	GDPMLISCLM	EHLYTEKMVE	DCEHRLLELQ	YFISRDWKLD	PVLYRKQCGD	ASRLCHTHGW	NETSEFMPOG	AVFSCLYRHA
			501	RALNEACESV	IQTACKHIRS	GDPMLISCLM	EHLYTEKMVE	DCEHRLLELQ	YFISRDWKLD	PVLYRKQCGD	ASRLCHTHGW	NETSEFMPOG	AVFSCLYRHA
			501	RALNEACESV	IQTACKHIRS	GDPMLISCLM	EHLYTEKMVE	DCEHRLLELQ	YFISRDWKLD	PVLYRKQCGD	ASRLCHTHGW	NETSEFMPOG	AVFSCLYRHA
23132	CFR-1	MG160	601	YRTEEQGRRL	SRECRAEVQR	ILHQRAMDVK	LDPALQDKCL	IDLGKWCSEK	TETGQLECL	QDHLDDLAVE	CRDIVGNLITE	LESEDIQIEA	ILMRACEPII
			601	YRTEEQGRRL	SRECRAEVQR	ILHQRAMDVK	LDPALQDKCL	IDLGKWCSEK	TETGQLECL	QDHLDDLAVE	CRDIVGNLITE	LESEDIQIEA	ILMRACEPII
			601	YRTEEQGRRL	SRECRAEVQR	ILHQRAMDVK	LDPALQDKCL	IDLGKWCSEK	TETGQLECL	QDHLDDLAVE	CRDIVGNLITE	LESEDIQIEA	ILMRACEPII
23132	CFR-1	MG160	701	QNFCHDVAON	QIDSGDLMEC	LIONKHQKDM	NEKCAIGVTH	FOLVQMKDFR	FSYKFMACK	EDVLKCLPNI	KKKVDVVICL	STTVRNDTLQ	EAKEHRVSLK
			701	QNFCHDVAON	QIDSGDLMEC	LIONKHQKDM	NEKCAIGVTH	FOLVQMKDFR	FSYKFMACK	EDVLKCLPNI	KKKVDVVICL	STTVRNDTLQ	EAKEHRVSLK
			701	QNFCHDVAON	QIDSGDLMEC	LIONKHQKDM	NEKCAIGVTH	FOLVQMKDFR	FSYKFMACK	EDVLKCLPNI	KKKVDVVICL	STTVRNDTLQ	EAKEHRVSLK
23132	CFR-1	MG160	801	CRRLRVEEL	EMTEDIRLEP	DLYEACKSDI	KNFCSAVQYG	NAQIIECLKE	NKKQLSTRCH	QKVEKLOETE	MMDPELDYTL	MRVCKQMIKK	FOPEADSKTM
			801	CRRLRVEEL	EMTEDIRLEP	DLYEACKSDI	KNFCSAVQYG	NAQIIECLKE	NKKQLSTRCH	QKVEKLOETE	MMDPELDYTL	MRVCKQMIKK	FOPEADSKTM
			801	CRRLRVEEL	EMTEDIRLEP	DLYEACKSDI	KNFCSAVQYG	NAQIIECLKE	NKKQLSTRCH	QKVEKLOETE	MMDPELDYTL	MRVCKQMIKK	FOPEADSKTM
23132	CFR-1	MG160	901	LQCLKQNKNS	ELMDPKCKQM	ITKRQITQNT	DYRLNPMRLK	ACKADIPKFC	HGILTKAKDD	SELEGQVISC	LKLRYADQRL	SSDCEDQIRI	IIQESALDYR
			901	LQCLKQNKNS	ELMDPKCKQM	ITKRQITQNT	DYRLNPMRLK	ACKADIPKFC	HGILTKAKDD	SELEGQVISC	LKLRYADQRL	SSDCEDQIRI	IIQESALDYR
			901	LQCLKQNKNS	ELMDPKCKQM	ITKRQITQNT	DYRLNPMRLK	ACKADIPKFC	HGILTKAKDD	SELEGQVISC	LKLRYADQRL	SSDCEDQIRI	IIQESALDYR
23132	CFR-1	MG160	1001	LDPQLQLHCS	DEISSLCAEE	AAAQEQTGQV	EECLKNLLK	IKTELCKKEV	LNMLKESKAD	IFVDPVLHTA	CALDIKHCA	AITPGRGRQM	SCLMEALEDK
			1001	LDPQLQLHCS	DEISSLCAEE	AAAQEQTGQV	EECLKNLLK	IKTELCKKEV	LNMLKESKAD	IFVDPVLHTA	CALDIKHCA	AITPGRGRQM	SCLMEALEDK
			1001	LDPQLQLHCS	DEISSLCAEE	AAAQEQTGQV	EECLKNLLK	IKTELCKKEV	LNMLKESKAD	IFVDPVLHTA	CALDIKHCA	AITPGRGRQM	SCLMEALEDK
23132	CFR-1	MG160	1101	RVRLQPECKK	RLNDRIEMWS	YAAKVAPADG	FSDLAMQVMT	SPSKNYILSV	ISGSICILFL	IGLMCGRITK	RVTRRELKDR*		
			1101	RVRLQPECKK	RLNDRIEMWS	YAAKVAPADG	FSDLAMQVMT	SPSKNYILSV	ISGSICILFL	IGLMCGRITK	RVTRRELKDR*		
			1101	RVRLQPECKK	RLNDRIEMWS	YAAKVAPADG	FSDLAMQVMT	SPSKNYILSV	ISGSICILFL	IGLMCGRITK	RVTRRELKDR*		

Fig. 16